

PCT_EP_04_00030_sequence listing.txt
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<110> alcedo biotech GmbH

<120> Use of HMGB, HMGN, HMGA proteins

<130> A 10009 PCT

<160> 64

<170> PatentIn version 3.1

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<213> Homo sapiens

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Ser Glu Val Pro Thr Pro Lys Arg Pro Arg Gly Arg Pro Lys Gly Ser
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Lys Asn Lys Gly Ala Ala Lys Thr Arg Lys Thr Thr Thr Pro Gly
65 70 75 80

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<213> Homo sapiens

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Arg Pro Lys Gly Ser Lys Asn Lys Gly Ala Ala Lys Thr Arg Lys Thr
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Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
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PCT_EP_04_00030_sequence listing.txt

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

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35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
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35 40 45

Trp Lys Thr Met Ser Ala Lys Glu Lys Gly Lys Phe Glu Asp Met Ala
50 55 60

Lys Ala Asp Lys Ala Arg Tyr Glu Arg Glu Met Lys Thr Tyr Ile Pro
65 70 75 80

Pro Lys Gly Glu Thr Lys Lys Phe Lys Asp Pro Asn Ala Pro Lys
85 90 95

Arg Pro Pro Ser Ala Phe Phe Leu Phe Cys Ser Glu Tyr Arg Pro Lys
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Leu Gly Glu Met Trp Asn Asn Thr Ala Ala Asp Asp Lys Gln Pro Tyr
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Glu Lys Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Ile Ala
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
65 70 75 80

Arg Lys Trp Ala Gly Val Gln Trp Tyr Asn Leu Gly Ser Leu Gln Pro
85 90 95

Pro Pro Pro Arg Phe Lys Gln Phe Ser Cys Leu Arg Leu Leu Ser Ser
100 105 110

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35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
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Arg Lys Trp Pro Gln Gln Val Val Gln Lys Lys Pro Ala Gln Tyr Ser
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

PCT_EP_04_00030_sequence listing.txt

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
65 70 75 80

Arg Lys Trp Pro Gln Gln Val Val Gln Lys Lys Pro Ala Gln Val Asn
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Phe Ser Lys Asn Ala Thr
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Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
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Phe Ser Glu Phe Ser Lys Lys Cys Ser Glu Arg Trp Lys Thr Met Ser
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Ala Lys Glu Lys Gly Lys Phe Glu Asp Met Ala Lys Ala Asp Lys Ala
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Arg Tyr Glu Arg Glu Met Lys Thr Tyr Ile Pro Pro Lys Gly

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Pro Arg Gly Lys Met Ser Ser Tyr Ala Phe Phe Val Gln Thr Cys Arg
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PCT_EP_04_00030_sequence listing.txt
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Glu Glu His Lys Lys Lys His Pro Asp Ala Ser Val Asn Phe Ser Glu
20 25 30

Phe Ser Lys Lys Cys Ser Glu Arg Trp Lys Thr Met Ser Ala Lys Glu
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Arg Glu Met Lys Thr Tyr Ile
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Pro Lys Lys Pro Arg Gly Lys Met Ser Ser Tyr Ala Phe Phe Val Gln
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Phe Ser Glu Phe Ser Lys Lys Cys Ser Glu Arg Trp Lys Thr Met Ser
35 40 45

Ala Lys Glu Lys Gly Lys Phe Glu Asp Met Ala Lys Ala Asp Lys Ala
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Arg Tyr Glu Arg Glu Met Lys Thr Tyr
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PCT_EP_04_00030_sequence listing.txt

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Glu Tyr Arg Pro Lys Ile Lys Gly Glu His Pro Gly Leu Ser Ile Gly
20 25 30

Asp Val Ala Lys Lys Leu Gly Glu Met Trp Asn Asn Thr Ala Ala Asp
35 40 45

Asp Lys Gln Pro Tyr Glu Lys Lys Ala Ala Lys Leu Lys Glu Lys Tyr
50 55 60

Glu Lys Asp Ile Ala Ala Tyr Arg Ala Lys Gly
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Pro Lys Arg Pro Pro Ser Ala Phe Phe Leu Phe Cys Ser Glu Tyr Arg
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Lys Lys Leu Gly Glu Met Trp Asn Asn Thr Ala Ala Asp Asp Lys Gln
35 40 45

Pro Tyr Glu Lys Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp
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Ile Ala Ala Tyr Arg
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PCT_EP_04_00030_sequence listing.txt

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Pro Lys Ile Lys Gly Glu His Pro Gly Leu Ser Ile Gly Asp Val Ala
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Lys Lys Leu Gly Glu Met Trp Asn Asn Thr Ala Ala Asp Asp Lys Gln
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Pro

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Glu Glu His Lys Lys Lys Asn Pro Asp Ala Ser Val Lys Phe Ser Glu
1 5 10 15

Phe Leu Lys Lys Cys Ser Glu Thr Trp Lys Thr Ile Phe Ala Lys Glu
20 25 30

Lys Gly Lys Phe Glu Asp Met Ala Lys Ala Asp Lys Ala His Tyr Glu
35 40 45

Arg Glu Met Lys Thr Tyr Ile Pro Pro Lys Gly Glu Lys Lys Lys Lys
50 55 60

Phe Lys Asp Pro Asn Ala Pro Lys Arg Pro Pro Leu Ala Phe Phe Leu
65 70 75 80

Phe Cys Ser Glu Tyr Arg Pro Lys Ile Lys Gly Glu His Pro Gly Leu
85 90 95

Ser Ile Asp Asp Val Val Lys Lys Leu Ala Gly Met Trp Asn Asn Thr
100 105 110

Ala Ala Ala Asp Lys Gln Phe Tyr Glu Lys Lys Ala Ala Lys Leu Lys
115 120 125

Glu Lys Tyr Lys Lys Asp Ile Ala Ala Tyr Arg Ala Lys Gly Lys Pro
130 135 140

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Glu Asp Asp Asp Lys
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20 25 30

Arg Lys Gln Gln Gln Glu Pro Thr Gly Glu Pro Ser Pro Lys Arg Pro
35 40 45

Arg Gly Arg Pro Lys Gly Ser Lys Asn Lys Ser Pro Ser Lys Ala Ala
50 55 60

Gln Lys Lys Ala Glu Ala Thr Gly Glu Lys Arg Pro Arg Gly Arg Pro
65 70 75 80

Arg Lys Trp Asn Thr Leu Glu Gln Cys Asn Val Cys Ser Lys Pro Ile
85 90 95

Met Glu Arg Ile Leu Arg Ala Thr Gly Lys Ala Tyr His Pro His Cys
100 105 110

Phe Thr Cys Val Met Cys His Arg Ser Leu Asp Gly Ile Pro Phe Thr
115 120 125

Val Asp Ala Gly Gly Leu Ile His Cys Ile Glu Asp Phe His Lys Lys
130 135 140

Phe Ala Pro Arg Cys Ser Val Cys Lys Glu Pro Ile Met Pro Ala Pro
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PCT_EP_04_00030_sequence listing.txt

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<210> 35

<211> 4111

<212> DNA

<213> Homo sapiens

<400> 35

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PCT_EP_04_00030_sequence listing.txt

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gtaagcaa	aa	at	at	tg	cc	a	ct	tt	tt	tt	tt	tt	1920
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<211> 330

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<213> Homo sapiens

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ggtagccct ctcctaagag acccaggggaa agacccaaag gcagcaaaaa caagagtccc 180

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tcctcacaag agtctgccga agaggactag 330

<210> 37

<211> 252

<212> DNA

<213> Homo sapiens

<400> 37

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ggtagccct ctcctaagag acccaggggaa agacccaaag gcagcaaaaa caagagtccc 180

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aggaaatggt ga 252

<210> 38

<211> 273

<212> DNA

PCT_EP_04_00030_sequence listing.txt

<213> Homo sapiens

<400> 38

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ggtagccct ctcctaagag acccagggga agacccaaag gcagcaaaaa caagagtccc
tctaaagcag ctcaaaaagaa agcagaagcc actggagaaaa aacggccaag aggccagacct
aggaaatggg aggagttta cattgcagct tag

60

120

180

240

273

<210> 39

<211> 291

<212> DNA

<213> Homo sapiens

<400> 39

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ggtagccct ctcctaagag acccagggga agacccaaag gcagcaaaaa caagagtccc
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291

<210> 40

<211> 1207

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> NCIB Accession No. NM_002128

<400> 40

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120

180

PCT_EP_04_00030_sequence listing.txt

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caagaggcct ccttcggcct tcttcctctt ctgctctgag tatcgcccaa aaatcaaagg	420
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<210> 41
<211> 648
<212> DNA
<213> Homo sapiens

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gccttcttcc tcttcgttc tgatgtatgc ccaaaaatca aaggagaaca tcctggcctg	360
tccattggtg atgttgcgaa gaaactggaa gagatgttggaa ataacactgc tgcagatgac	420
aagcagcctt atgaaaagaa ggctgcgaag ctgaaggaaa aatacgaaaa ggatattgt	480

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gcatatcgag ctaaaggaaa gcctgatgca gcaaaaaagg gagttgtcaa ggctaaaaaa 540
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gaagatgaag aagatgaaga tgaagaagaa gatgatgatg atgaataa 648

<210> 42

<211> 444

<212> DNA

<213> Homo sapiens

<400> 42
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ggtgagccct ctcctaagag acccaggggaa agacccaaag gcagcaaaaa caagagtccc 180
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cacccagcta atttttgtat ttttagtaga gacagggttt caccatgttgc gccaggctgg 420
tctcgaactc ctgacacctcag gtga 444

<210> 43

<211> 321

<212> DNA

<213> Homo sapiens

<400> 43
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ggtgagccct ctcctaagag acccaggggaa agacccaaag gcagcaaaaa caagagtccc 180
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aggaaatggg acaatctact accaagaacc agctccaaga agaaaacatc tctggaaac 300
agtaccaaaa ggagtcaactg a 321

<210> 44

<211> 279

PCT_EP_04_00030_sequence listing.txt

<212> DNA

<213> Homo sapiens

<400> 44

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ggtgagccct ctcctaagag acccagggga agacccaaag gcagcaaaaa caagagtccc 180
tctaaagcag ctcaaaagaa agcagaagcc actggagaaa aacggccaag aggtagac 240
aggaaatggt ggttgctaattt gaagagcccg tgctggtaa 279

<210> 45

<211> 291

<212> DNA

<213> Homo sapiens

<400> 45

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aggaaatggc cacaacaagt tgttcagaag aagcctgctc agtattcctg a 291

<210> 46

<211> 357

<212> DNA

<213> Homo sapiens

<400> 46

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aggaaatggc cacaacaagt tgttcagaag aagcctgctc aggtcaatgt tgccttgcc 300
gggaaggacc acccggccaa tcttatatat ctactgttct ctaaaaatgc cacttag 357

PCT_EP_04_00030_sequence listing.txt

<210> 47

<211> 288

<212> DNA

<213> Homo sapiens

<400> 47

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ggtgagccct ctcctaagag acccagggga agacccaaag gcagcaaaaa caagagtccc 120
tctaaagcag ctcaaaagaa agcagaagcc actggagaaa aacggccaag aggtagac 180
aggaaaatggc cacaacaagt tgttcagaag aagcctgctc aggactga 240
288

<210> 48

<211> 33

<212> DNA

<213> Homo sapiens

<400> 48

actgagaagc ggggccccggg caggccgcgc aag 33

<210> 49

<211> 33

<212> DNA

<213> Homo sapiens

<400> 49

acacctaaga gacctgggg ccgaccaaag gga 33

<210> 50

<211> 36

<212> DNA

<213> Homo sapiens

<400> 50

PCT_EP_04_00030_sequence listing.txt
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36

<210> 51
<211> 33
<212> DNA
<213> Homo sapiens

<400> 51
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33

<210> 52
<211> 33
<212> DNA
<213> Homo sapiens

<400> 52
acacctaaga gacctcgggg ccgaccaaag gga

33

<210> 53
<211> 36
<212> DNA
<213> Homo sapiens

<400> 53
actccaggaa ggaaaccaag gggcagaccc aaaaaa

36

<210> 54
<211> 33
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<213> Homo sapiens

<400> 54
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33

<210> 55
<211> 33

PCT_EP_04_00030_sequence listing.txt

<212> DNA

<213> Homo sapiens

<400> 55
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<210> 56

<211> 63

<212> DNA

<213> Homo sapiens

<400> 56
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aag 63

<210> 57

<211> 234

<212> DNA

<213> Homo sapiens

<400> 57
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tcagagaggt ggaaggtaag agggcttaaa acatgctaac aaggtaatta aaagacagtt 180
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<210> 58

<211> 213

<212> DNA

<213> Homo sapiens

<400> 58
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tggaagacca tgtctgctaa agagaaagga aaatttgaag atatggcaaa ggcggacaag 180

PCT_EP_04_00030_sequence listing.txt

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<210> 59

<211> 219

<212> DNA

<213> Homo sapiens

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tcagagaggt ggaagaccat gtctgctaaa gagaaaggaa aatttgaaga tatggcaaag 180
gcggacaagg cccgttatga aagagaaaatg aaaacctat 219

<210> 60

<211> 225

<212> DNA

<213> Homo sapiens

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aaggaaaaat acgaaaagga tattgctgca tatcgagcta aagga 225

<210> 61

<211> 207

<212> DNA

<213> Homo sapiens

<400> 61
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aacactgctg cagatgacaa gcagccttat gaaaagaagg ctgcgaagct gaaggaaaa 180
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PCT_EP_04_00030_sequence listing.txt

<210> 62

<211> 147

<212> DNA

<213> Homo sapiens

<400> 62
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aacactgctg cagatgacaa gcagcct 147

<210> 63

<211> 546

<212> DNA

<213> Homo sapiens

<400> 63
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aaataa 546

<210> 64

<211> 678

<212> DNA

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